
Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: Fri Oct 12 13:51:26 EDT 2007

Validated By CRFValidator v 1.0.3

Application No: 10561235 Version No: 2.0

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Finished: 2007-10-01 17:39:32.948

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 354 ms

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Total Errors: 0

No. of SeqIDs Defined: 7

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215

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Phe Glu	Ala Ph	e Lys	Gln	Tyr	His 120	Leu	Asp	Leu	Leu	Gln 125	Asn	Ser	Glu
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Arg Trp Ala Gly Ala Val Leu Phe Ala Ala Ala Phe Val Ser Val Arg 130 135 140

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Asp	Ala	Pro	Asp	Leu 405	Asn	Ala	Lys	Thr	Val 410	Ser	Ala	Leu	Asn	Asp 415	Tyr
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Val Gly	Lys Ser	Met Thr 85	Lys Ala	Pro Met 90	Ile Asp	Phe Ser	Val Val 95
Ser Arg	Asn Gly	Val Ala	Ala Leu	Ala Gly 105	Asp Gln	Tyr Ile 110	Val Ser
Val Ala	His Asn	Gly Gly	Tyr Asn 120	Asn Val	Asp Phe	Gly Ala 125	Glu Gly
Ser Asn 130	Pro Asp	Gln His	Arg Phe	Ser Tyr	Gln Ile 140	Val Lys	Arg Asn
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Met Pro	Arg Leu	His Lys 165	Phe Val	Thr Asp	Ala Glu	Pro Val	Glu Met 175
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Thr Gly As	n Pro Tyr	lle Gly 295	_	Ser Asn	Gly Ph		Leu	Val	Arg
Lys Asp Tr 305	p Phe Tyr	Asp Glu	Ile E	Phe Ala	Gly As	p Thr	His	Ser	Val 320
Phe Tyr Gl	u Pro His 325		Gly I	Lys Tyr 330	Phe Ph	ie Asn	Asp	Asn 335	Asn
Asn Gly Al	a Gly Lys 340	lle Asp		Lys His 345	Lys Hi	s Tyr	Ser 350	Leu	Pro
Tyr Arg Le	_	Arg Thr	Val (Gln Leu	Phe As	n Val 365	Ser	Leu	Ser
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Tyr Arg Pr 385	o Arg Leu	Asn Asn 390	Gly (Glu Asn	Ile Se	er Phe	Ile	Asp	Lys 400
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Trp Lys Va 450	l Asn Gly	Val Ala 455		Asp Arg	Leu Se		Ile	Gly	Lys
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